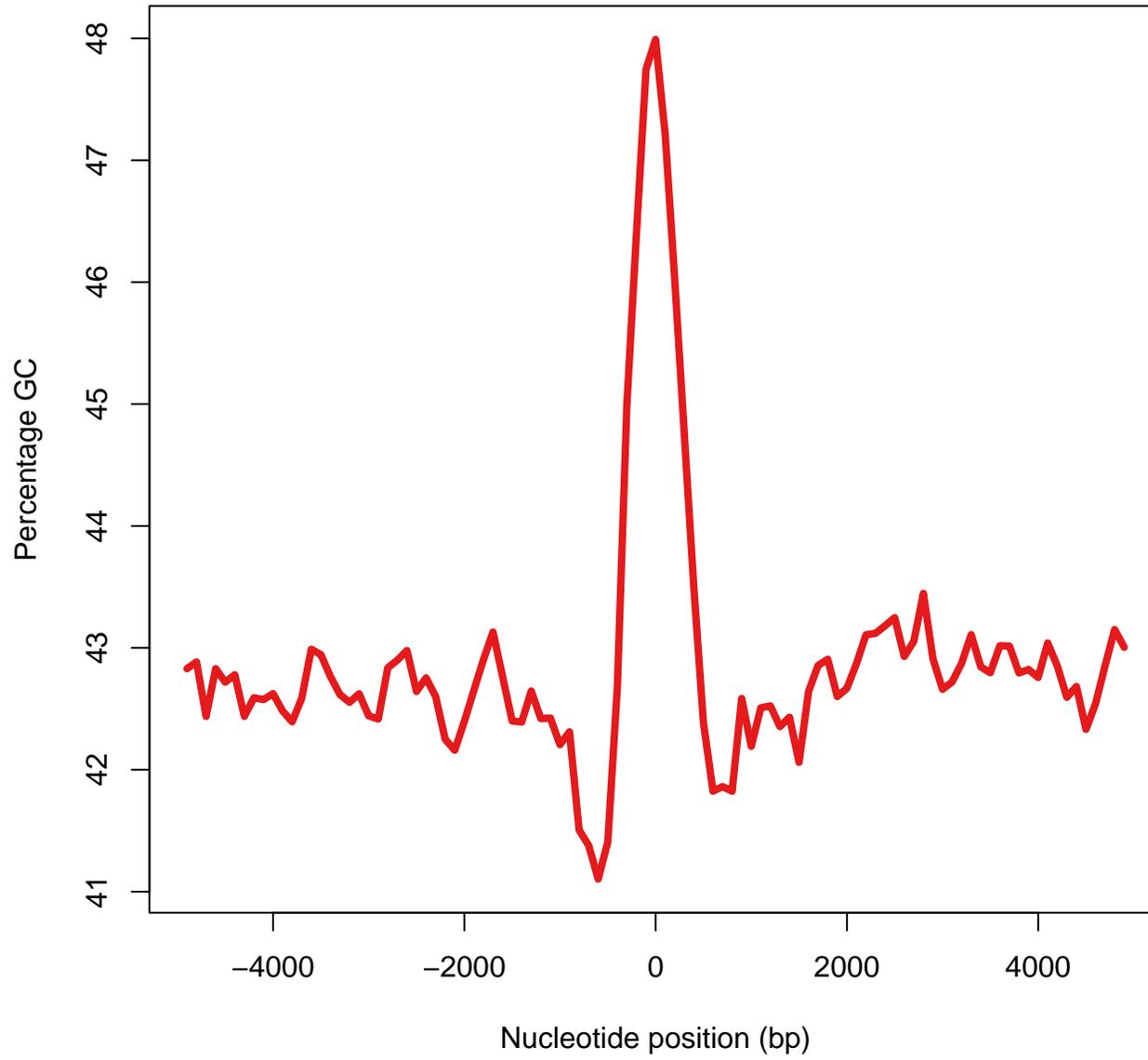
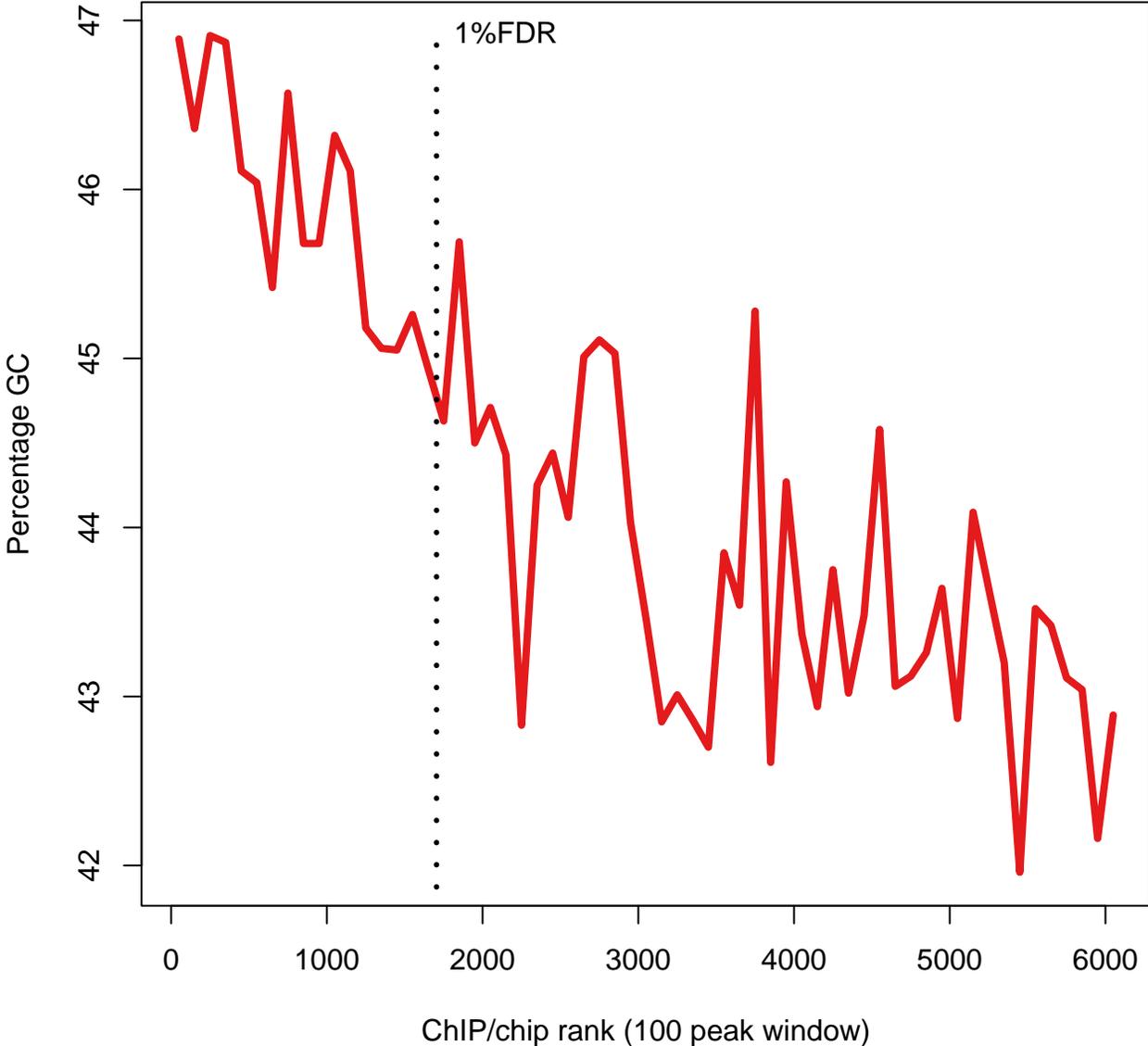


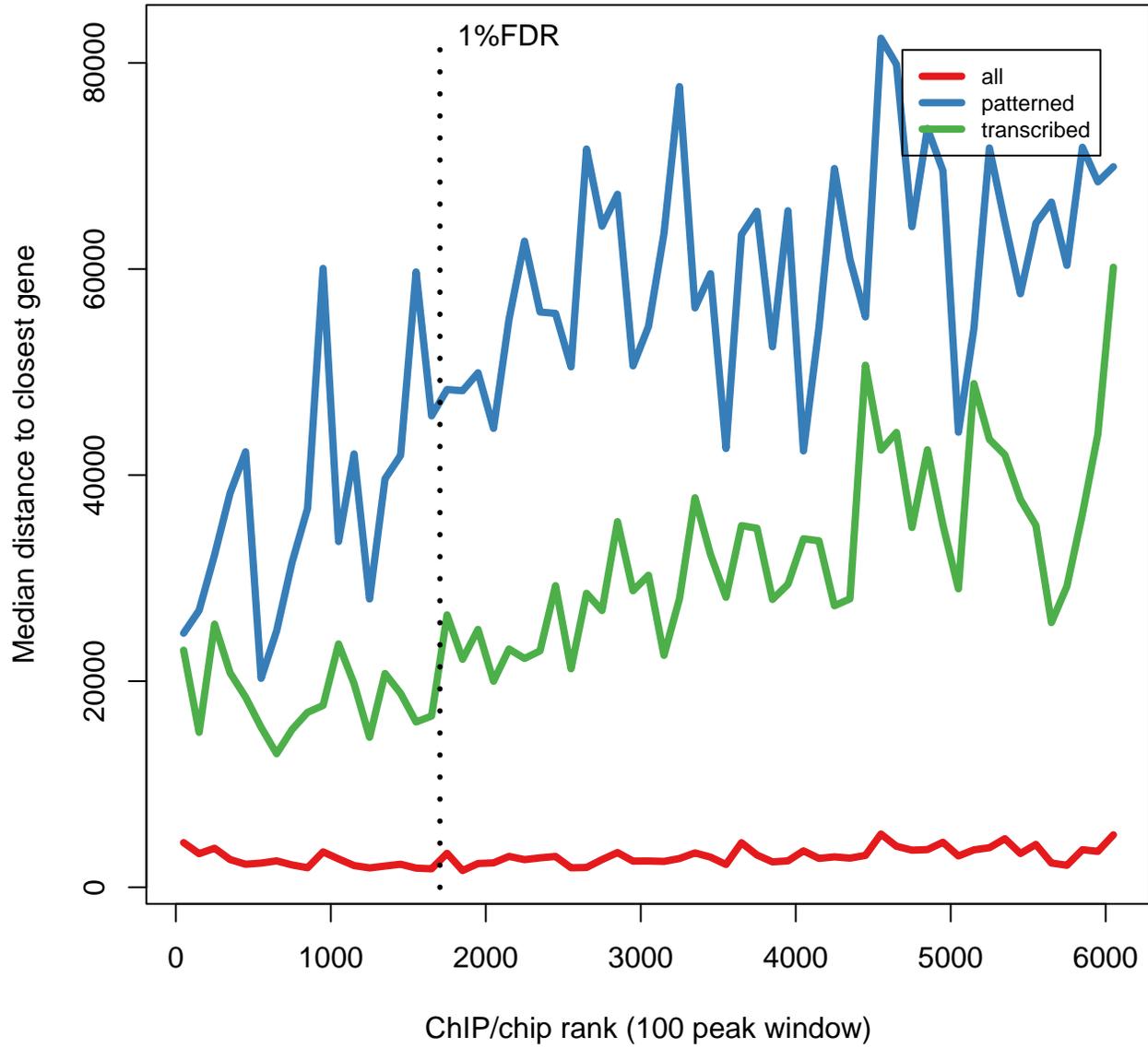
**h\_1 base composition across peaks**



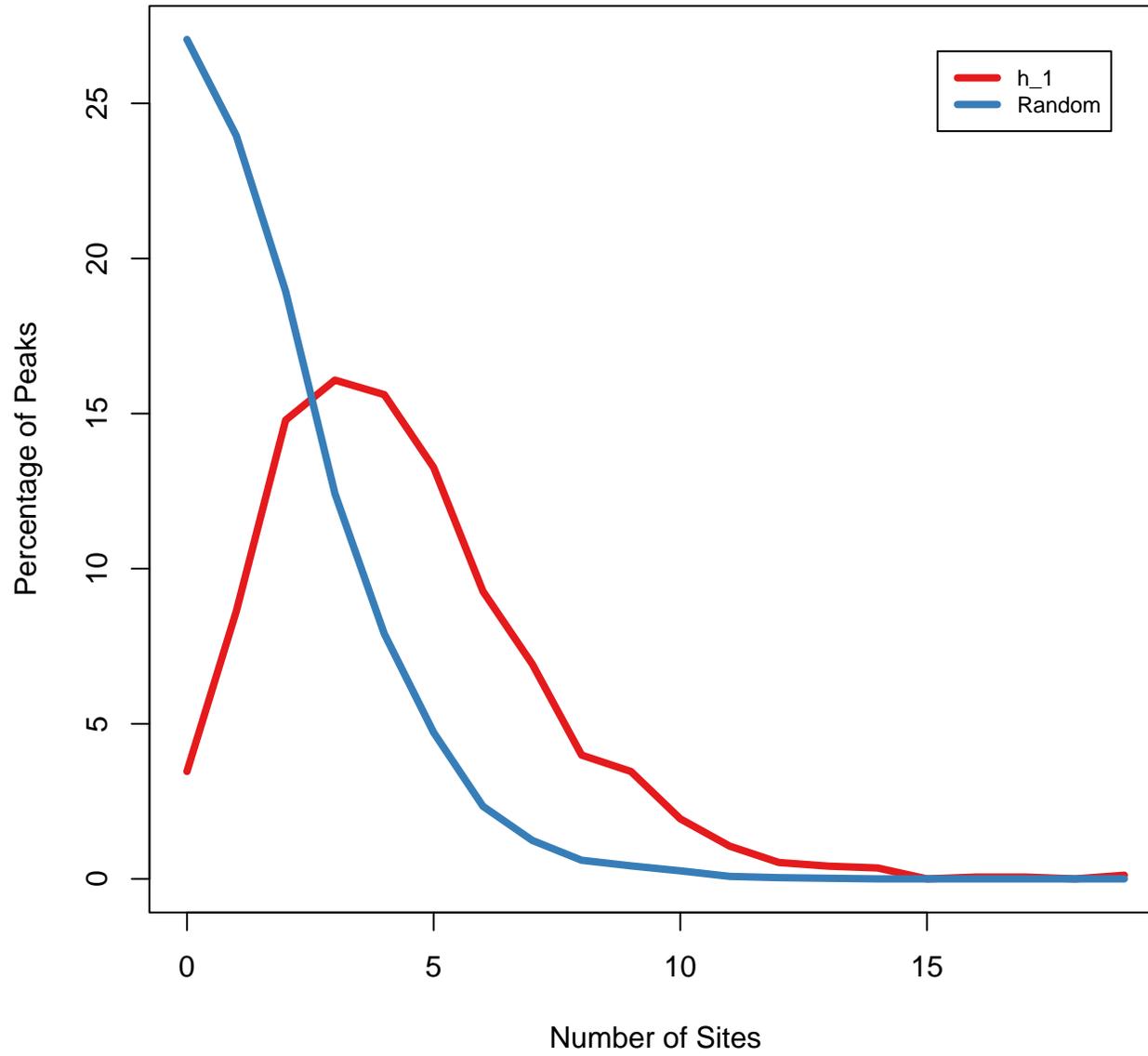
**h\_1 base composition down ranks**



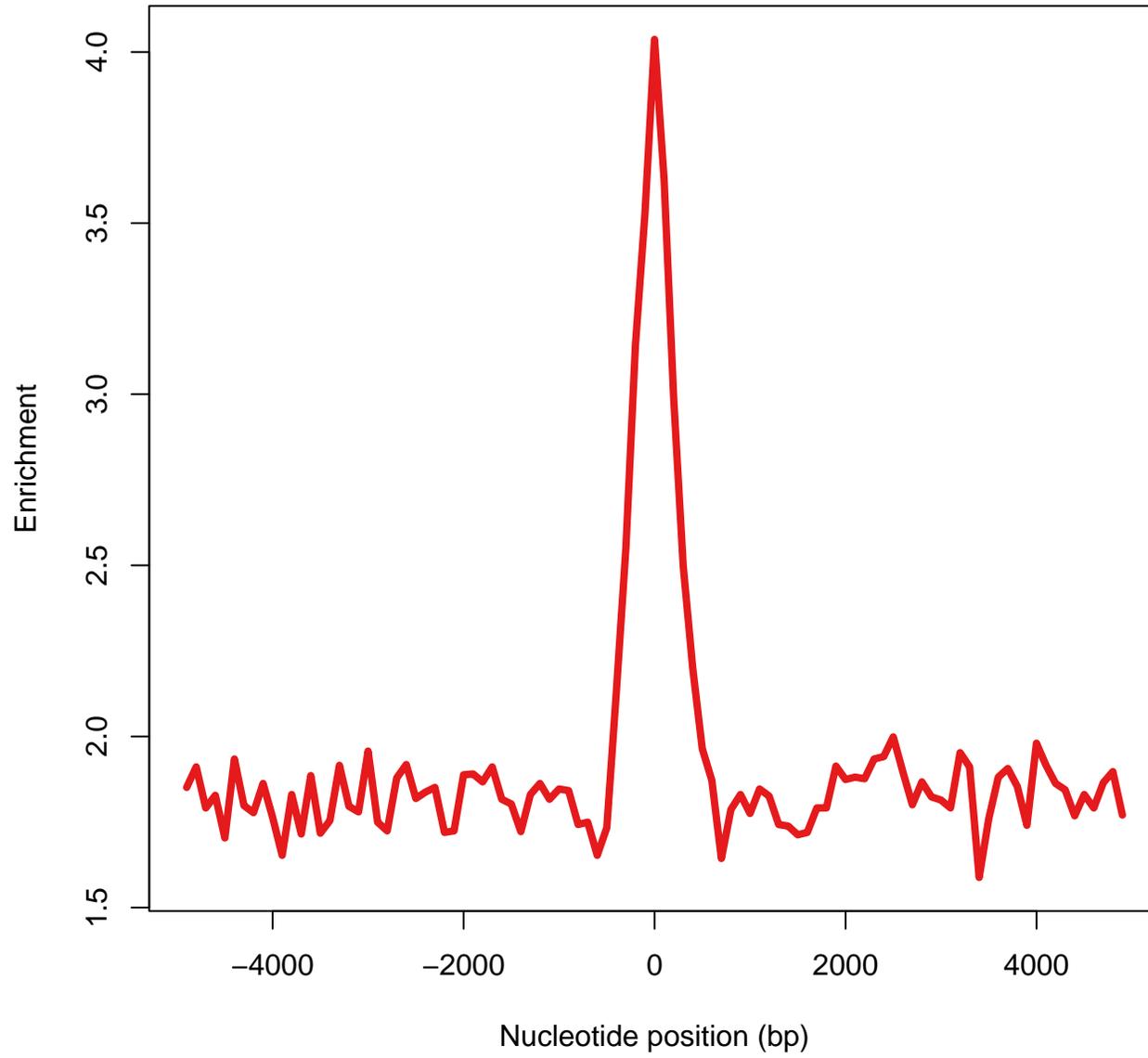
### h\_1 median distance to genes



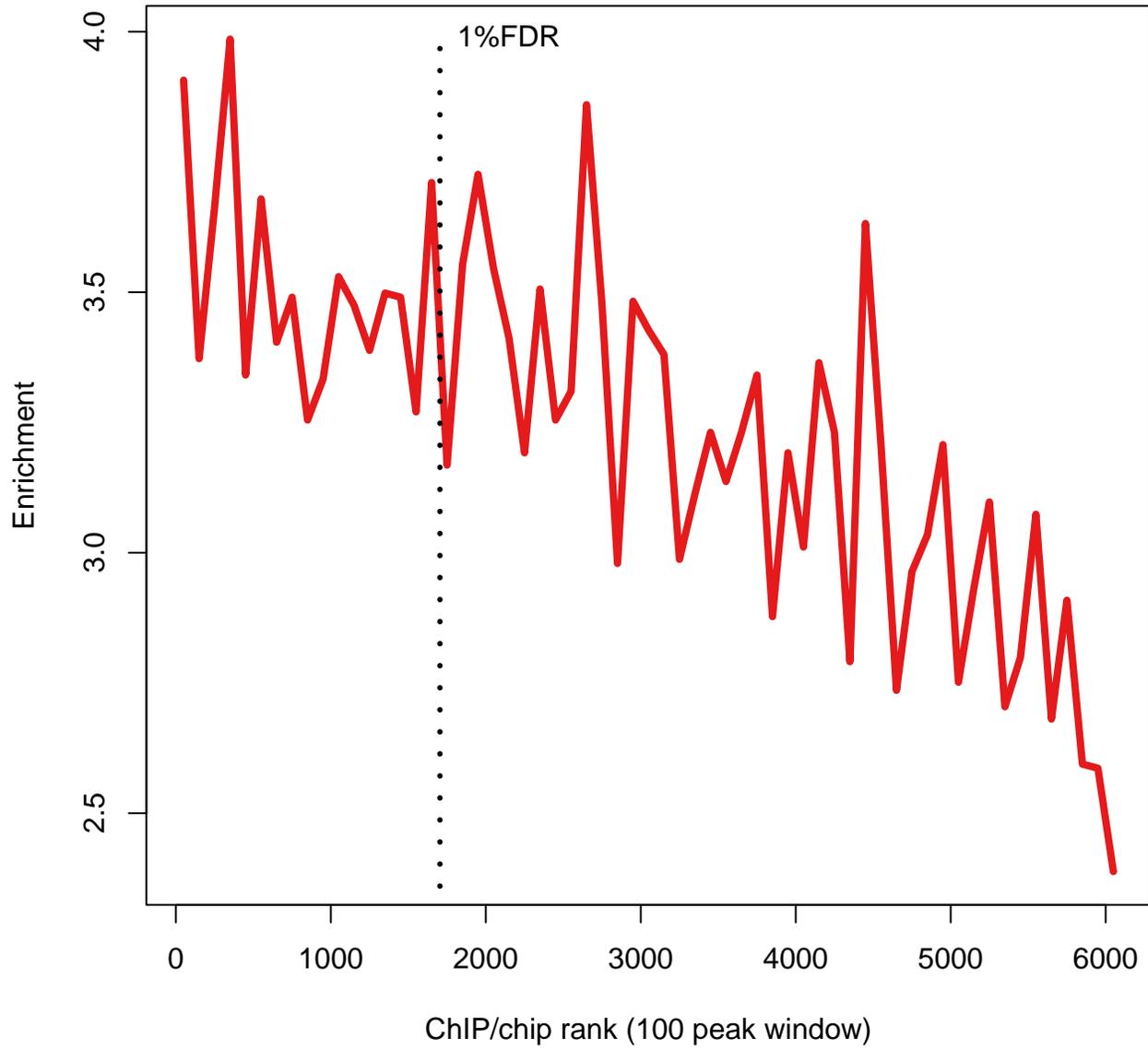
**h<sub>1</sub> distribution of number of sites**



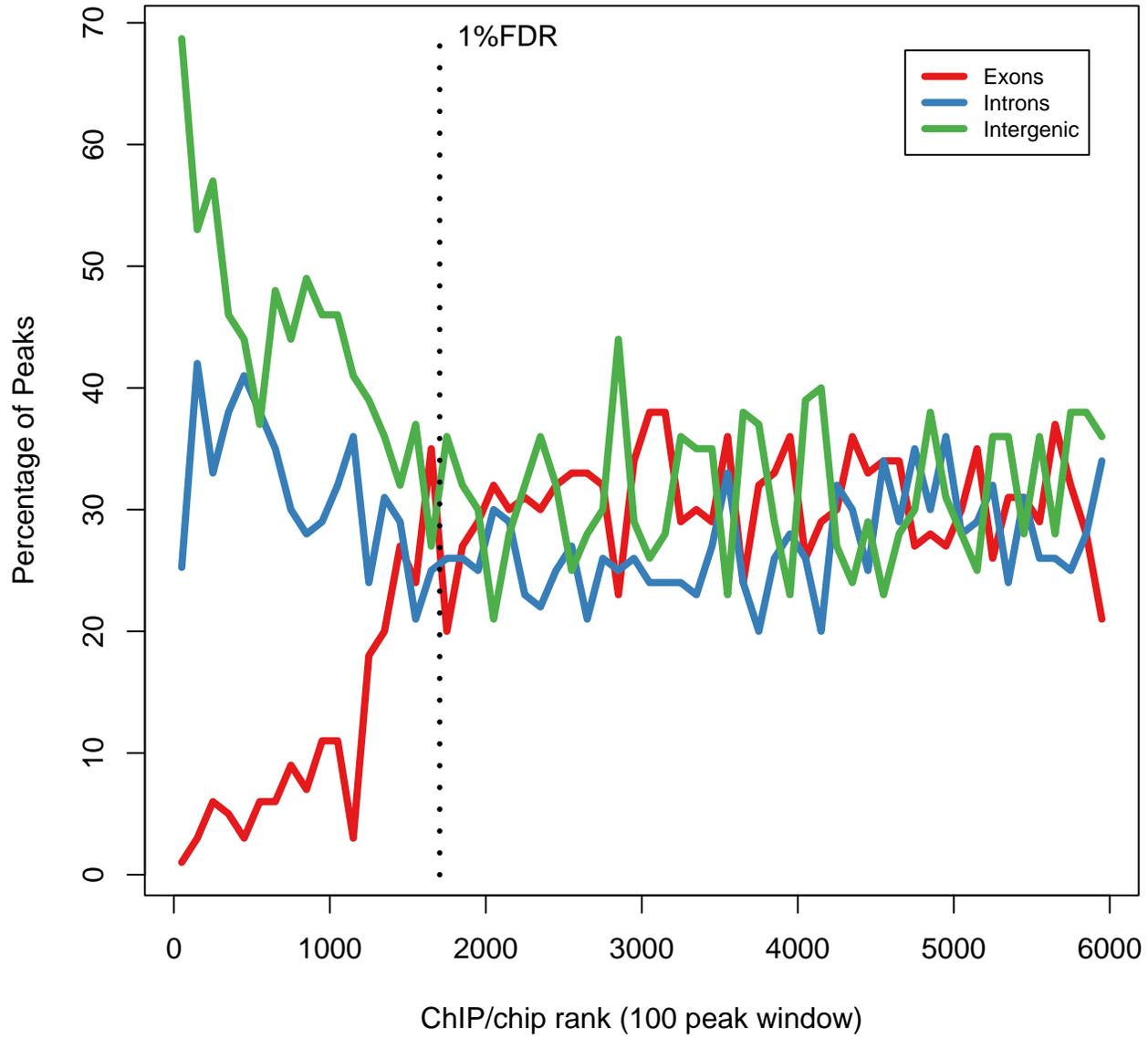
**h\_1 PWM enrichment across peaks**



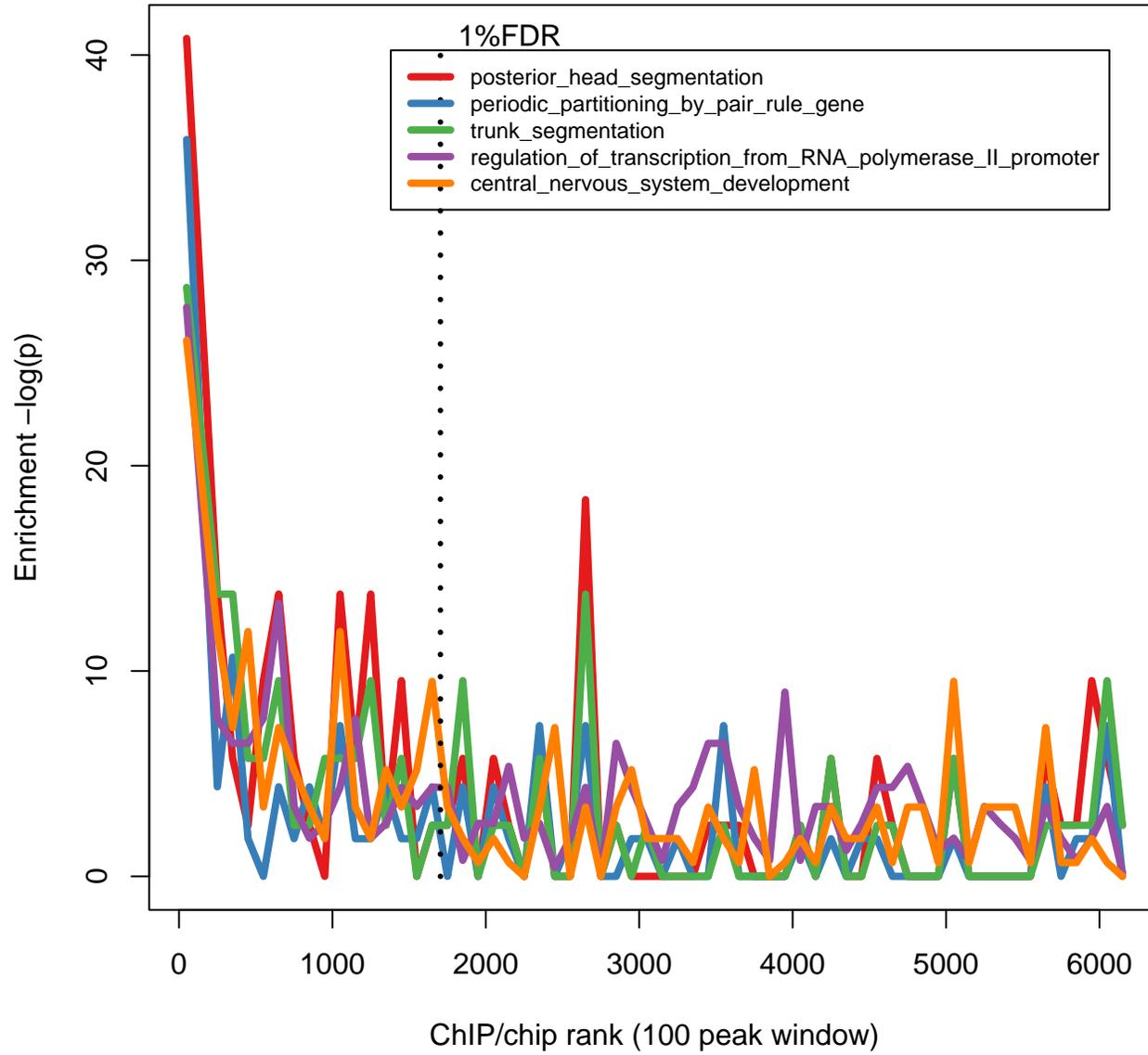
### h\_1 PWM enrichment down ranks

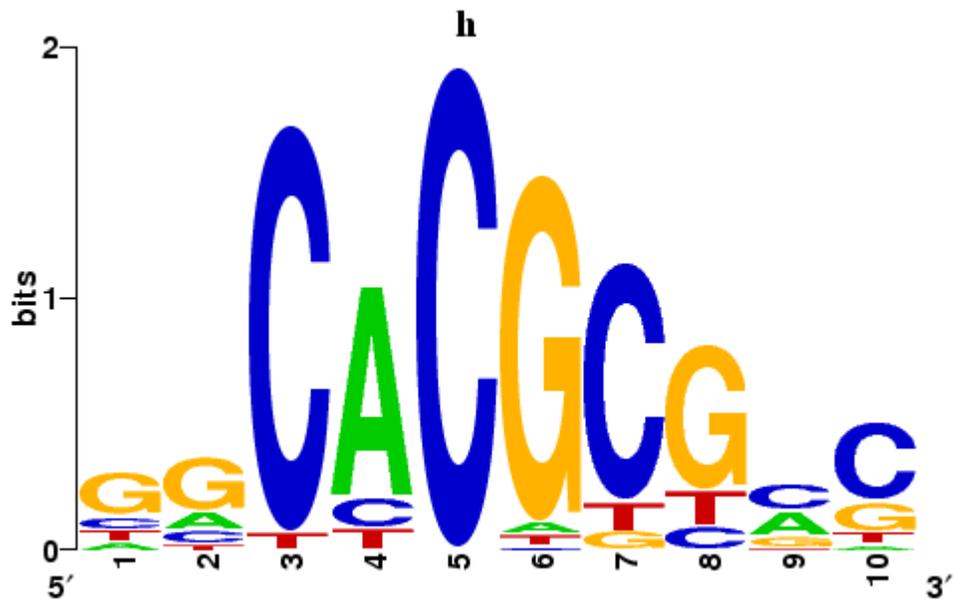


# h\_1 genomic location

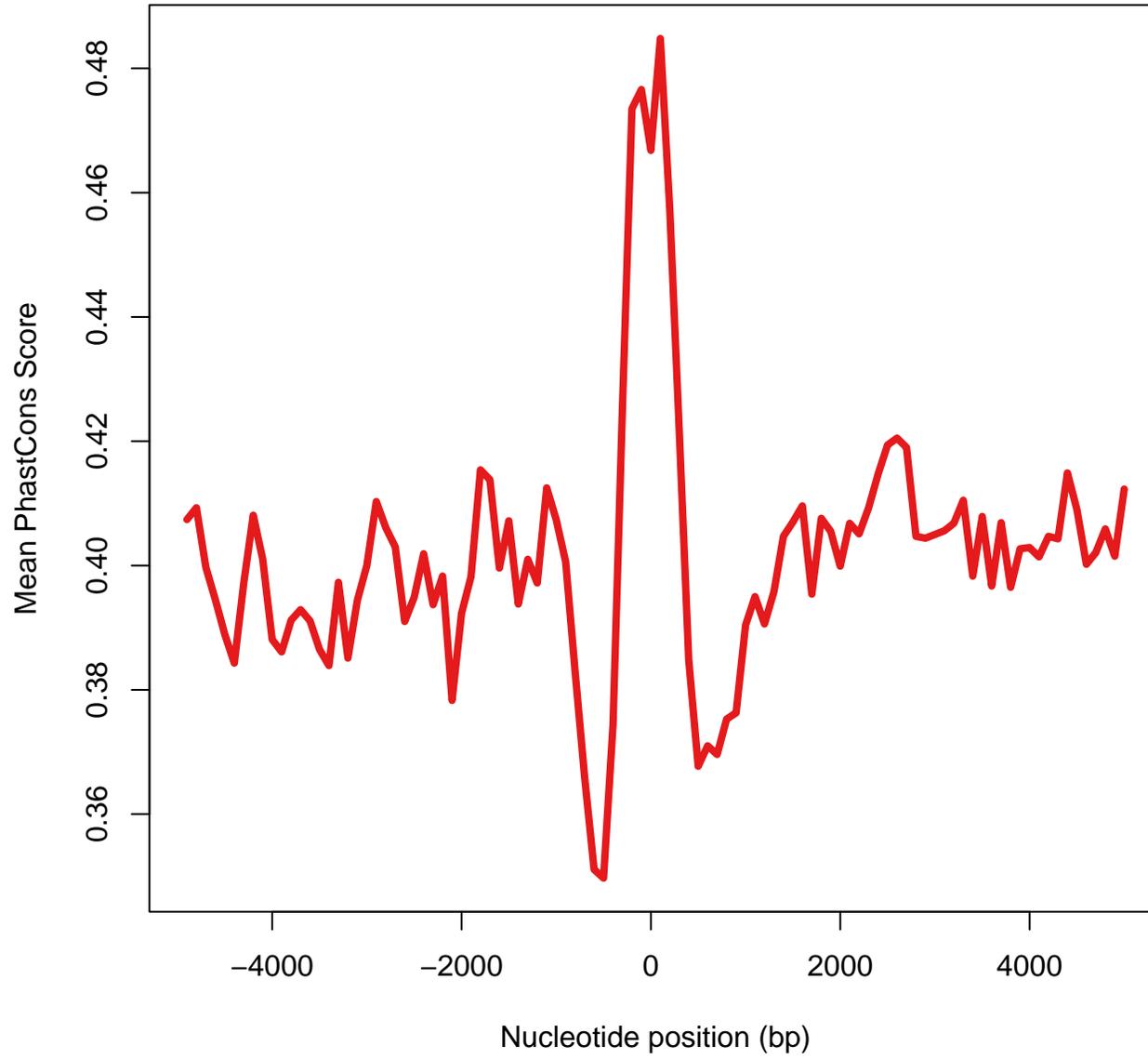


### h\_1 GO term enrichment (-logP)





**h\_1 mean PhastCons across peaks**



**h\_1 mean PhastCons down ranks**

